

Page 1 of 7

1655

RAW SEQUENCE LISTING DATE: 02/27/2001
PATENT APPLICATION: US/09/698,341 TIME: 15:27:52

Input Set : A:\sequencelist.txt

Output Set: N:\CRF3\02272001\1698341.raw

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3 <110> APPLICANT: Sorge, Joseph
         Hurlbut Hogrefe, Holly
         Connie, Hansen
 7 <120> TITLE OF INVENTION: Compositions and Methods Utilizing DNA Polymerases
                                                                                  ENTERED
 9 <130> FILE REFERENCE: 25436/1560
11 <140> CURRENT APPLICATION NUMBER: 09/698,341
12 <141> CURRENT FILING DATE: 2000-10-27
14 <150> PRIOR APPLICATION NUMBER: 60/162,600
15 <151> PRIOR FILING DATE: 1999-10-29
17 <160> NUMBER OF SEQ ID NOS: 48
19 <170> SOFTWARE: PatentIn version 3.0
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22 <211> LENGTH: 2331
23 <212> TYPE: DNA
24 <213> ORGANISM: Thermococcus sp. JDF-3
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31 etecteaggg acgaetetge categaagaa ateaaaaaga taacegegga gaggeacgge
33 agggtegtta aggttaageg egeggagaag gtgaagaaaa agtteetegg eaggtetgtg
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35 gaggtotggg tectotactt caegeaeceg caggaegtte eggeaateeg egacaaaata
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37 aggaagcacc ccgcggtcat cgacatctac gagtacgaca tacccttcgc caagcgctac
39 cteatagaca agggeetaat ecogatggaa ggtgaggaag agettaaaet catyteette
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41 gacatcgaga cgctctacca cgagggagaa gagtttggaa ccgggccgat tctgatgata
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43 agetaegeeg atgaaagega ggegegegtg ataacetgga agaagatega cetteettae
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45 gttgaggttg tctccaccga gaaggagatg attaagcgct tcttgagggt cgttaaggag
                                                                          600
47 aaggaccegg acgtgctgat aacatacaac ggcgacaact tcgacttcgc ctacctgaaa
49 aagogetgtg agaagettgg egtgagettt acceteggga gggaegggag egageegaag
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51 atacagegea tgggggacag gtttgeggte gaggtgaagg gcagggtaca ettegacett
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53 tatecagtea taaggegeae cataaacete eegacetaca eeettgagge tgtataegag
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55 geggtttteg geaageecaa ggagaaggte taegeegagg agatageeae egeetgggag
57 acceggegagg ggcttgagag ggtcgcgcgc tactcgatgg aggacgcgag ggttacctac
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{\tt 59}\ {\tt gagettggca}\ {\tt gggagttctt}\ {\tt cccgatggag}\ {\tt geccagettt}\ {\tt ccaggetcat}\ {\tt cggccaagge}
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61 ctctgggacg tttcccgctc cagcaccggc aacctcgtcg agtggttcct cctaaggaag
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63 gcctacgaga ggaacgaact cgctcccaac aagcccgacg agagggaget ggcgaggaga
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65 aggggggget acgccggtgg ctacgtcaag gagccygagc ggggactgtg ggacaatatc
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67 gtgtatotag aetttegtag tetotaccet teaateataa teaeceacaa egtetegeea
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69 gatacgctca accycgaggg gtgtaggagc tacgacgttg cccccgaggt cggtcacaag
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71 ttctgcaagg acttccccgg cttcattccg agcctgctcg gaaacctgct ggaggaaagg
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73 cagaagataa agaggaagat gaaggcaact etegaceege tggagaagaa teteetegat
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83 atggagttot taaactatat caateecaaa etgeeeggee ttetegaact egaataegag
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85 ggcttctacg tcaggggctt cttcgtcacg aagaaaaagt acgcggtcat cgacgaggag
87 ggcaagataa ccacgegegg gettgagata gtcaggegeg actggagega gatagegaag
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Input Set : A:\sequencelist.txt
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89 gagacgcagg cgagggtttt ggaggcgata ctcaggcacg gtgacgttga agaggccgtc 1920 91 agaattgtca gggaagtcac cgaaaagetg agcaagtacg aggtteegec ggagaagetg 1980 93 gttatecaeg ageagataac gegegagete aaggactaca aggeeaeegg eeegeaegta 95 gocatagoga agogtttgge ogecagaggt gttaaaatce ggcooggaac tgtgataage 2100 97 tacatogtto tgaagggote oggaaggata ggogacaggg ogattooott ogacgagtto 2160 99 gaccegacga agcacaagta cgatgeggac tactacateg agaaccaggt tetgeeggea 2220 101 gttgagagaa teeteaggge etteggetae egeaaggaag acetgegeta eeagaagaeg 103 aggcaggtcg ggcttggcgc gtggctgaag ccgaagggga agaagaagtg a 106 <210> SEQ ID NO: 2 107 <211> LENGTH: 776 108 <212> TYPE: PRT 109 <213> ORGANISM: Thermococcus sp. JDF-3 111 <400> SEQUENCE: 2 113 Met Ile Leu Asp Val Asp Tyr Ile Thr Glu Asn Gly Lys Pro Val Ile 10 116 Arg Val Phe Lys Lys Glu Asn Gly Glu Phe Arg Ile Glu Tyr Asp Arg 117 20 25 30 119 Glu Phe Glu Pro Tyr Phe Tyr Ala Leu Leu Arg Asp Asp Ser Ala Ile 120 35 40 122 Glu Glu Ile Lys Lys Ile Thr Ala Glu Arg His Gly Arg Val Val Lys 123 50 55 60 125 Val Lys Arg Ala Glu Lys Val Lys Lys Lys Phe Leu Gly Arg Ser Val 126 65 70 70 80 128 Glu Val Trp Val Leu Tyr Phe Thr His Pro Gln Asp Val Pro Ala Ile 129  $$85\ 90\ 95$ 131 Arg Asp Lys Ile Arg Lys His Pro Ala Val Ile Asp Ile Tyr Glu Tyr 132  $\phantom{\bigg|}100\phantom{\bigg|}105\phantom{\bigg|}105\phantom{\bigg|}$ 134 Asp Ile Pro Phe Ala Lys Arg Tyr Leu Ile Asp Lys Gly Leu Ile Pro 135  $$1.5\,$  L15  $$120\,$ 137 Met Glu Glu Glu Glu Leu Lys Leu Met Ser Phe Asp Ile Glu Thr 138 130 135 140 140 Leu Tyr His Glu Gly Glu Glu Phe Gly Thr Gly Pro Ile Leu Met Ile 141 145 150 150 160 143 Ser Tyr Ala Asp Glu Ser Glu Ala Arg Val Ile Thr Trp Lys Lys Ile 144  $\phantom{\bigg|}$  165  $\phantom{\bigg|}$  170  $\phantom{\bigg|}$  170  $\phantom{\bigg|}$  175 146 Asp Leu Pro Tyr Val Glu Val Val Ser Thr Glu Lys Glu Met Ile Lys 147  $180 \hspace{1.5cm} 185 \hspace{1.5cm} 185 \hspace{1.5cm} 190$ 149 Arg Phe Leu Arg Val Val Lys Glu Lys Asp Pro Asp Val Leu Ile Thr 150 195 200 205 152 Tyr Asn Gly Asp Asn Phe Asp Phe Ala Tyr Leu Lys Lys Arg Cys Glu 153 210 215 220 155 Lys Leu Gly Val Ser Phe Thr Leu Gly Arg Asp Gly Ser Glu Pro Lys 156 225  $230 \hspace{1.5cm} 230 \hspace{1.5cm} 235 \hspace{1.5cm} 235$ 158 lie Gln Arg Met Gly Asp Arg Phe Ala Val Glu Val Lys Gly Arg Val 159 245 250 255 161 His Phe Asp Leu Tyr Pro Val Ile Arg Arg Thr Ile Asn Leu Pro Thr 162 260 265 270 164 Tyr Thr Leu Glu Ala Val Tyr Glu Ala Val Phe Gly Lys Pro Lys Glu 280

TECH CENTER 1600/290

RAW SEQUENCE LISTING

PATENT APPLICATION: US/09/698,341

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167 Lys Val Tyr Ala Glu Glu Ile Ala Thr Ala Trp Glu Thr Gly Glu Gly 168 290 295 300 170 Leu Glu Arg Val Ala Arg Tyr Ser Met Glu Asp Ala Arg Val Thr Tyr 171 305 310 315 173 Glu Leu Gly Arg Glu Phe Phe Pro Met Glu Ala Gln Leu Ser Arg Leu 174  $\phantom{\bigg|}325\phantom{\bigg|}335\phantom{\bigg|}$  330  $\phantom{\bigg|}335\phantom{\bigg|}$ 176 Tle Gly Gln Gly Leu Trp Asp Val Ser Arg Ser Ser Thr Gly Asn Leu 177  $\phantom{\bigg|}340\phantom{\bigg|}$  345  $\phantom{\bigg|}350\phantom{\bigg|}$ 179 Val Glu Trp Phe Leu Leu Arg Lys Ala Tyr Glu Arg Asn Glu Leu Ala 180  $\phantom{\bigg|}355\phantom{\bigg|}360\phantom{\bigg|}360\phantom{\bigg|}365\phantom{\bigg|}$ 182 Pro Asn Lys Pro Asp Glu Arg Glu Leu Ala Arg Arg Arg Gly Gly Tyr 183  $\phantom{+}370\phantom{+}375\phantom{+}375\phantom{+}380\phantom{+}$ 185 Ala Gly Gly Tyr Val Lys Glu Pro Glu Arg Gly Leu Trp Asp Asn Ile 186 385  $390 \hspace{1.5cm} 395 \hspace{1.5cm} 395$ 188 Val Tyr Leu Asp Phe Arg Ser Leu Tyr Pro Ser Ile Ile Ile Thr  $\operatorname{His}$ 405 410 191 Asıı Val Ser Pro Asp Thr Leu Asıı Arg Glu Gly Cys Arg Ser Tyr Asp 192 420 425 430 194 Val Ala Pro Glu Val Gly His Lys Phe Cys Lys Asp Phe Pro Gly Phe 195  $\phantom{\bigg|}435\phantom{\bigg|}435\phantom{\bigg|}440\phantom{\bigg|}$ 197 The Pro Ser Leu Leu Gly Asn Leu Leu Glu Glu Arg Gln Lys Ile Lys 198  $\phantom{\bigg|}450\phantom{\bigg|}455\phantom{\bigg|}$ 200 Arg Lys Met Lys Ala Thr Leu Asp Pro Leu Glu Lys Asn Leu Leu Asp 201 465 470 475 203 Tyr Arg Gln Arg Ala Ile Lys Tle Leu Ala Asn Ser Tyr Tyr Gly Tyr 204 485 490 495 206 Tyr Gly Tyr Ala Arg Ala Arg Trp Tyr Cys Arg Glu Cys Ala Glu Ser 207  $\phantom{\bigg|}500\phantom{\bigg|}505\phantom{\bigg|}510\phantom{\bigg|}$ 209 Val Thr Ala Trp Gly Arg Glu Tyr Ile Glu Met Val Ile Arg Glu Leu 210 515 520 525 212 Glu Glu Lys Phe Gly Phe Lys Val Leu Tyr Ala Asp Thr Asp Gly Leu 213  $\phantom{\bigg|}530\phantom{\bigg|}535\phantom{\bigg|}535\phantom{\bigg|}540\phantom{\bigg|}$ 215 His Ala Thr Ile Pro Gly Ala Asp Ala Glu Thr Val Lys Lys Lys Ala 216 545 550 560 218 Met Glu Phe Leu Asn Tyr Ile Asn Pro Lys Leu Pro Gly Leu Leu Glu 219  $\phantom{\bigg|}565\phantom{\bigg|}570\phantom{\bigg|}570\phantom{\bigg|}575\phantom{\bigg|}$ 224 Lys Tyr Ala Val Ile Asp Glu Glu Gly Lys Ile Thr Thr Arg Gly Leu 225 595 600 605 · 227 Glu Ile Val Arg Arg Asp Trp Ser Glu Ile Ala Lys Glu Thr Gln Ala 228  $\,$  610  $\,$  615  $\,$  620 233 Arg Ile Val Arg Glu Val Thr Glu Lys Leu Ser Lys Tyr Glu Val Pro 234 645 650 655 650 655 236 Pro Glu Lys Leu Val fle His Glu Gln Ile Thr Arg Glu Leu Lys Asp 237 660 665 670 239 Tyr Lys Ala Thr Gly Pro His Val Ala Ile Ala Lys Arg Leu Ala Ala

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242 Arg Cly Val Lys 1le Arg Pro Cly Thr Val Ile Ser Tyr Ile Val Leu
243 690 695 700
245 Lys Gly Ser Gly Arg 11e Gly Asp Arg Ala Ile Pro Phe Asp Glu Phe 246 705 710 715 720
248 Asp Pro Thr Lys His Lys Tyr Asp Ala Asp Tyr Tyr Ile Glu Asp Gln 249 725 \cdot \phantom{1}735
251 Val Leu Pro Ala Val Glu Arg Tle Leu Arg Ala Phe Gly Tyr Arg Lys 252 740 745 750
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279 Glu Phe Glu Pro Tyr Phe Tyr Ala Leu Leu Arg Asp Asp Ser Ala Ile 280 35 40 45
282 Glu Glu Ile Lys Lys Ile Thr Ala Glu Arg His Gly Arg Val Val Lys 283 \phantom{000}55\phantom{000} 50
285 Val Lys Arg Ala Glu Lys Val Lys Lys Lys Phe Leu Gly Arg Ser Val 286 65 70 75 80
288 Glu Val Trp Val Leu Tyr Phe Thr His Pro Gln Asp Val Pro Ala Ile
289 85 90 95
291 Arg Asp Lys Ile Arg Lys His Pro Ala Val Ile Asp Ile Tyr Glu Tyr 292 \phantom{\bigg|}100\phantom{\bigg|}100\phantom{\bigg|}105\phantom{\bigg|}
294 Asp Ile Pro Phe Ala Lys Arg Tyr Leu Ile Asp Lys Gly Leu Ile Pro 295 \phantom{\bigg|}115\phantom{\bigg|}120\phantom{\bigg|}120\phantom{\bigg|}125\phantom{\bigg|}
297 Met Glu Glu Glu Glu Glu Leu Lys Leu Met Ser Phe Asp Ile Glu Thr 298 \phantom{\bigg|}130 \phantom{\bigg|}135 \phantom{\bigg|}135
300 Leu Tyr His Glu Gly Glu Glu Phe Gly Thr Gly Pro Ile Leu Met Ile 301 145 \phantom{\bigg|} 150 \phantom{\bigg|} 150 \phantom{\bigg|} 155 \phantom{\bigg|} 160
303 Ser Tyr Ala Asp Glu Ser Glu Ala Arg Val Ile Thr Trp Lys Lys Ile 304 \phantom{\bigg|}165\phantom{\bigg|}170\phantom{\bigg|}170\phantom{\bigg|}170\phantom{\bigg|}
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309 Arg Phe Leu Arg Val Val Lys Glu Lys Asp Pro Asp Val Leu Ile Thr
310 195
                           200
312 Tyr Asn Gly Asp Asn Phe Asp Phe Ala Tyr Leu Lys Lys Arg Cys Glu
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RAW SEQUENCE LISTING .

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	225		2			230					235					240
	Tle	Gln	Arq	Met	Glv		Arg	Phe	Ala	Val	Clu	Val	Lys	Cly	Arq	Val
319			5		245		5			250			2 -		255	
	His	Phe	Asp	Leu	Tyr	Pro	Val	He	Arq	Arg	Thr	Tle	Asn	Leu	Pro	Thr
322				260					265	_				270		
	Tyr	Thr	Leu	Glu	Ala	Val.	Tyr	Glu	Ala	Val	Phe	Gly	Lys	Pro	Lys	Glu
325	•		275				•	280				•	285		•	
327	Lys	Val	Tyr	Ala	Glu	Glu	I l.e	Ala	Thr	Ala	Trp	Glu	Thr	Gly	Glu	Gly
328	-	290	-				295				•	300		-		
330	Leu	Glu	Arg	Val	Ala	Arg	Туr	ser	Met	Glu	Asp	Al.a	Arg	Va1	Thr	Tyr
331	305		-			310	_				315					320
333	G.l.u	Leu	GLy	Arg	Glu	Phe	Phe	Pro	Met	G.l.u	Ala	Gln	Leu	Ser	Arg	Leu
334					325					330					335	
336	Ile	Gly	Gln	Gly	Leu	Trp	Asp	Val	Ser	Arg	Ser	ser	Thr	Gly	Asn	Leu
337				340					345					350		
339	Val	Glu	Trp	Phe	Leu	Leu	Arg	Lys	Ala	Tyr	$\operatorname{Glu}$	Arg	Asn	Glu	Leu	Ala
340			355					360					365			
342	Pro	Asn	Lys	Pro	Asp	Glu	Arg	Glu	Leu	Ala	Arg	Arg	Arg	Gly	Gly	Tyr
343		370					375					380				
345	Ala	Gly	Gly	Tyr	Val.	ràs	Glu	Pro	Glu	Arg		Leu	Trp	Asp	Asn	Ile
346	385					390					395					400
348	Val	Tyr	Leu	Asp		Arg	ser	Leu	Tyr		Ser	11e	Ile	Ile		His
349					405					410				•	415	
	Asn	Val	Ser		Asp	Thr	Leu	Asn		Glu	Gly	Cys	Arg		Tyr	Asp
352	_			420					425			_		430		
	Val	Ala		GLu	Va.L	Gly	His		Phe	Cys	Lys	Asp		Pro	GTA	Phe
355	~ 7		435		_			440		er 2	a		445	<b>.</b>	*1.	
	Ile		ser	ren	Leu	GIY		ren	Leu	GIU	GLU	-	GIN	rys	TTG	Lys
358	1	450	1104	T	2.7	ml	455		Dwa	T a	<i>c</i> 1	460	Nan	T 0	T	7.00
	Arg	riAz	Mec	гаг	ATA	470	Leu	ASP	PLO	ren	475	nys	ASII	Leu	ren	480
361	465 Tyr	7 210	Cln	V ~ ~	7. 1 ->		Lug	110	T ou	7.15		Car	Lou	T Ou	Dro	
364	тут	Arg	GIII	MIG	485	TTE	пЛр	116	цеи	490	MSII	361	пен	Leu	495	СТУ
	G.l u	Trn	Val	Δla		Tle	Glu	Glv	Glv		T.e.u	Arα	Pro	Va 1		TTe
367	0.1 4	11.12	¥ C2 3.	500	ru.L	1.1.0	OLU	0.17	505	ny o		**** 9	0	510	**** 9	1. 1. 0
	Gly	Glu	Len		Asp	Gly	Len	Met		Ala	Ser	Glv	Glu		Val	Lvs
370	OLY	O.L.	515	V CL 1.	, rot	017	1100	520	O I II	riiu	001	(11)	525	1119	,	1,5
	Arg	Asp		Asp	Thr	Glu	Val		Glu	Val	Glu	Glv		Tvr	Ala	Ser
373	**** 9	530	0.27			-	535					540		-1-		
	Pro		Thr	Gly	ser	Pro		Lvs	Pro	Ala	Gln	Cvs	Arg	Lvs	Pro	Glv
376	545			-		550	-	•			555	•	-	-		560
378	Thr	Ala	Met	P.ro	G1y	Lys	Phe	Thr	Glu	Leu	Ser	Thr	Pro	Glu	Gly	Gly
379					565	-				570					575	•
381	Leu	ser	Val	Thr	Arg	Gly	His	Ser	Leu	Phe	Al.a	Tyr	Arg	Asp	Ala	Ser
382				580		_			585			-	-	590		
384	Leu	Trp	A.rg	Arg	Gly	Arg	Arg	Arg	Phe	Lys	Pro	Gly	Asp	Leu	Leu	Ala
385			595					600					605			



## Please Note:

Use of n and/or Xaa have been detected in the Sequence Listing. Please review the Sequence Listing to ensure that a corresponding explanation is presented in the <220> to <223> fields of each sequence which presents at least one n or Xaa.



VERIFICATION SUMMARY

PATENT APPLICATION: US/09/698,341

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Input Set : A:\sequencelist.txt

Output Set: N:\CRF3\02272001\1698341.raw

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L:656 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:4
L:690 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:4
L:692 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:4
L:736 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:5
L:758 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:6
L:774 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:7
L:970 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:21
L:1146 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:22
L:1235 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:28
L:1275 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:29
L:1340 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:31
L:1380 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:32
L:1420 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:33
L:1739 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:33
L:1739 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:33